Amendments to the Claims

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims

- (Currently Amended) An isolated nucleic acid encoding an M. catarrhalis polypeptide of <u>SEQ ID</u>
 NO:3218 SEQ ID NOS: 1921 [[-]] 3840.
- 2. (Original) A recombinant expression vector comprising the nucleic acid of Claim 1 operably linked to a transcription regulatory element.
- 3. (Original) A cell comprising a recombinant expression vector of Claim 2.
- 4. (Original) A method for producing an *M. catarrhalis* polypeptide comprising culturing a cell of Claim 3 under conditions that permit expression of the polypeptide.
- 5. (Currently Amended) An isolated nucleic acid selected from the group consisting of:
 - (a) <u>SEQ ID NO:1298 SEQ ID NOS: 1</u> [[-]] 1920;
 - (b) a complement of <u>SEQ ID NO:1298</u> SEQ ID NOS: 1 [[-]] 1920; or
 - (c) an RNA of (a) or (b), wherein U is substituted for T.
- 6. (Original) A recombinant expression vector comprising the nucleic acid of Claim 5 operably linked to a transcription regulatory element.
- 7. (Original) A cell comprising a recombinant expression vector of Claim 6.
- 8. (Original) A method for producing an *M. catarrhalis* polypeptide comprising culturing a cell of Claim 7 under conditions that permit expression of the polypeptide.
- 9. (Currently Amended) A probe comprising a nucleotide sequence consisting of at least eight contiguous nucleotides of a nucleotide sequence selected from the group consisting of:
 - (a) <u>SEQ ID NO:1298</u> SEQ ID NOS: 1 [[-]] 1920;
 - (b) a complement of <u>SEQ ID NO:1298</u> SEQ ID NOS: 1 [[-]] 1920; or

- (c) an RNA of (a) or (b), wherein U is substituted for T.
- 10. (Currently Amended) An isolated nucleic acid comprising a nucleotide sequence of at least eight nucleotides in length, wherein the sequence is hybridizable to a nucleic acid having a nucleotide sequence selected from the group consisting of:
 - (a) <u>SEQ ID NO:1298</u> <u>SEQ ID NOS: 1</u> [[-]] <u>1920</u>;
 - (b) a complement of <u>SEQ ID NO:1298</u> SEQ ID NOS:1 [[-]] 1920; or
 - (c) an RNA of (a) or (b), wherein U is substituted for T.
- 11. (Withdrawn) A vaccine composition for prevention or treatment of an *M. catarrhalis* infection comprising a nucleic acid of Claim 5 and a pharmaceutically acceptable carrier.
- 12. (Withdrawn) A vaccine composition of Claim 11, further comprising an adjuvant.
- 13. (Withdrawn) A vaccine composition of Claim 11, further comprising one or more additional ingredients.
- 14. (Withdrawn) A method of treating a subject for *M. catarrhalis* infection comprising administering to a subject a vaccine composition of Claim 11, such that treatment of *M. catarrhalis* infection occurs.
- 15. (Withdrawn) A method of Claim 14, wherein the treatment is a prophylactic treatment.
- 16. (Withdrawn) A method of Claim 14, wherein the treatment is a therapeutic treatment.
- 17. (Withdrawn) A recombinant or substantially pure preparation of an *M. catarrhalis* polypeptide or a fragment thereof, wherein said *M. catarrhalis* polypeptide is SEQ ID NOS: 1921-3840.
- 18. (Withdrawn) A vaccine composition for prevention or treatment of an *M. catarrhalis* infection comprising an *M. catarrhalis* polypeptide of Claim 17 and a pharmaceutically acceptable carrier.
- 19. (Withdrawn) A vaccine composition of Claim 18, further comprising an adjuvant.

- (Withdrawn) A vaccine composition of Claim 18, further comprising one or more additional ingredients.
- 21. (Withdrawn) A method of treating a subject for *M. catarrhalis* infection comprising administering to a subject a vaccine composition of Claim 18, such that treatment of *M. catarrhalis* infection occurs.
- 22. (Withdrawn) A method of Claim 21, wherein the treatment is a prophylactic treatment.
- 23. (Withdrawn) A method of Claim 21, wherein the treatment is a therapeutic treatment.
- 24. (Withdrawn) A method for detecting the presence or absence of a *Klebsiella* nucleic acid in a sample comprising:
 - (a) contacting a sample with the nucleic acid of Claim 5 under conditions in which a hybrid can form between a probe comprising a nucleotide sequence consisting of at least eight contiguous nucleotides of a nucleotide sequence selected from the group consisting of SEQ ID NOS: 1-2501 or a complement of SEQ ID NOS: 1-1920 and a *Klebsiella* nucleic acid in the sample; and
 - (b) detecting the hybrid formed in step (a), wherein detection of a hybrid indicates the presence or absence of a *Klebsiella* nucleic acid in the sample.
- 25. (Withdrawn) A computer readable medium having recorded thereon a nucleotide sequence selected from the group consisting of:
 - (a) SEQ ID NOS: 1-1920;
 - (b) a complement of SEQ ID NOS: 1- 1920;
 - (c) an RNA of (a) or (b), wherein U is substituted for T; or
 - (d) a fragment of (a), (b) or (c).
- 26. (Withdrawn) A computer based system for identifying fragments of the *Klebsiella* genome of comprising;
 - a) a data storage means comprising a nucleotide sequence selected from the group consisting of SEQ ID NOS: 1-1920, a complement of SEQ ID NOS: 1-1920, or a fragment thereof,

- b) a search means for comparing a target sequence to the nucleotide sequences of the data storage means of step (a) to identify homologous sequences, and;
- c) a retrieval means for obtaining said homologous sequences(s) of step (b).
- 27. (Withdrawn) A method of identifying nucleic acid fragments of a Klebsiella genome comprising comparing a database comprising a nucleotide sequence selected from the group consisting of SEQ ID NOS: 1-1920; a complement of SEQ ID NOS: 1-1920; or a fragment thereof with a target sequence to obtain a nucleic acid molecule comprised of a complementary nucleotide sequence to said target sequence, wherein said target sequence is not randomly selected.
- 28 (Withdrawn) A method for identifying an expression modulating fragment of the *Klebsiella* genome comprising comparing a database comprising a nucleotide sequence selected from the group consisting of SEQ ID NOS: 1- 1920; a complement of SEQ ID NOS: 1-1920; or fragment thereof with a target sequence to obtain a nucleic acid molecule comprised of a complementary nucleotide sequence to said target sequence, wherein said target sequence comprises sequences known to regulate gene expression.